## SEQUENCE LISTING

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EULENBERG, KARSTEN
NGUYEN, TRI
BURK, ULRIKE
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- <120> USE OF A SECRETED PROTEIN FOR PREVENTING AND TREATING OBESITY, DIABETES AND/OR METABOLIC SYNDROME
- <130> 18744-0033
- <140> 10/550,985
- <141> 2005-09-26
- <150> PCT/EP04/003244
- <151> 2004-03-26
- <150> EP 03006948.8
- <151> 2003-03-26
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- <213> Homo sapiens

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<213> Homo sapiens

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Ile Cys Lys Asp Val Val Thr Ala Ala Gly Asp Met Leu Lys Asp Asn

Ala Thr Glu Glu Ile Leu Val Tyr Leu Glu Lys Thr Cys Asp Trp

Leu Pro Lys Pro Asn Met Ser Ala Ser Cys Lys Glu Ile Val Asp Ser .

Tyr Leu Pro Val Ile Leu Asp Ile Ile Lys Gly Glu Met Ser Arg Pro 120

Gly Glu Val Cys Ser Ala Leu Asn Leu Cys Glu Ser Leu Gln Lys His 130

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- Leu Leu Eu Tyr Pro Gln Asp Gly Pro Arg Ser Lys Pro Gln Pro Lys
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- Asp Asn Gly Asp Val Cys Gln Asp Cys Ile Gln Met Val Thr Asp Ile 195 200 205
- Gln Thr Ala Val Arg Thr Asn Ser Thr Phe Val Gln Ala Leu Val Glu 210 215 220
- His Val Lys Glu Glu Cys Asp Arg Leu Gly Pro Gly Met Ala Asp Ile 225 230 235 240
- Cys Lys Asn Tyr Ile Ser Gln Tyr Ser Glu Ile Ala Ile Gln Met Met 245 250 255
- Met His Met Gln Pro Lys Glu Ile Cys Ala Leu Val Gly Phe Cys Asp 260 265 270
- Glu Val Lys Glu Met Pro Met Gln Thr Leu Val Pro Ala Lys Val Ala 275 280 285
- Ser Lys Asn Val Ile Pro Ala Leu Glu Leu Val Glu Pro Ile Lys Lys 290 295 300
- His Glu Val Pro Ala Lys Ser Asp Val Tyr Cys Glu Val Cys Glu Phe 305 310 315 320
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- Glu Ile Leu Asp Ala Phe Asp Lys Met Cys Ser Lys Leu Pro Lys Ser 340 345 350
- Leu Ser Glu Glu Cys Gln Glu Val Val Asp Thr Tyr Gly Ser Ser Ile 355 360 365
- Leu Ser Ile Leu Leu Glu Glu Val Ser Pro Glu Leu Val Cys Ser Met 370 375 380
- Leu His Leu Cys Ser Gly Thr Arg Leu Pro Ala Leu Thr Val His Val 385 390 395 400
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- Gly Tyr Leu Asp Arg Asn Leu Glu Lys Asn Ser Thr Lys Gln Glu Ile 420 425 430
- Leu Ala Ala Leu Glu Lys Gly Cys Ser Phe Leu Pro Asp Pro Tyr Gln
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- Lys Gln Cys Asp Gln Phe Val Ala Glu Tyr Glu Pro Val Leu Ile Glu 450 455 460

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Cys Arg Ala Leu Gly His Cys Leu Gln Glu Val Trp Gly His Val Gly
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Asn	Gln	Val 115	Leu	Asp	Asp	Tyr	Phe 120	Pro	Leu	Val	Ile	Asp 125	Tyr	Phe	Glr
Asn	Gln 130	Ile	Asp	Ser	Asn	Gly 135	Ile	Cys	Met	His	Leu 140	Gly	Leu	Cys	Lys
Ser 145	Arg	Gln	Pro	Glu	Pro 150	Glu	Gln	Glu	Pro	Gly 155	Met	Ser	Asp	Pro	Leu 160
Pro	Lys	Pro	Leu	Arg 165	Asp	Pro	Leu	Pro	Asp 170	Pro	Leu	Leu	Asp	Lys 175	Leu
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Thr	Gln	Asp 195	Leu	Ser	Glu	Gln	Gln 200	Phe	Pro	Ile	Pro	Leu 205	Pro	Tyr	Суя
Trp	Leu 210	Cys	Arg	Ala	Leu	Ile 215	Lys	Arg	Ile	Gln	Ala 220	Met	Ile	Pro	Lys
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Gln Asp Ile Ala Ala Ala Gly Asn Gly Leu Asn Pro Asp Ala Thr

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70

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- Gln Val Cys Thr Ala Leu Ser Leu Cys Glu Pro Leu Gln Arg His Leu 130 135 140
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- Glu Gly Ala Leu Cys Gln Asp Cys Val Arg Gln Val Ser Arg Leu Gln 180 185 190
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Thr Val Ser Ser His Asn Leu Glu Ser Lys Ser Thr Lys Arg Asp Ile 405 410 415

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Ile Gln Cys Lys His Phe Val Thr Gln Tyr Glu Pro Val Leu Ile Glu 435 440 445

Ser Leu Lys Asp Met Met Asp Pro Val Ala Val Cys Lys Lys Val Gly 450 455 460

Ala Cys His Gly Pro Arg Thr Pro Leu Leu Gly Thr Asp Gln Cys Ala 465 470 475 480

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Gln Ala Arg Asp Gln Leu Lys Ser Asn Gln Thr Glu Glu Glu Leu Lys 85 90 95

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180

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- Pro Ile Asp His Leu Met Gly Ala Ala Asn Pro Gly Ala Leu Val Glu
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- Gly Gly Glu Leu Cys Thr Leu Cys Glu Tyr Met Leu His Phe Ile Gln 435 440 445

- Glu Thr Leu Ala Thr Pro Ser Thr Asp Asp Glu Ile Lys His Thr Val 450 455 460
- Glu Asn Ile Cys Ala Lys Leu Pro Ser Gly Val Ala Gly Gln Cys Arg 465 470 475 480
- Asn Phe Val Glu Met Tyr Gly Asp Ala Val Ile Ala Leu Leu Val Gln 485 490 495
- Gly Leu Asn Pro Arg Asp Val Cys Pro Leu Met Gln Met Cys Pro Lys
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- Asn Leu Pro Lys Lys Glu Asp Val Glu Val Phe Asn Pro Gln Pro Ala 515 520 525
- Ser Asp Glu Gln Asp Pro Pro Thr Cys Pro Leu Cys Leu Phe Ala Val 530 540
- Glu Gln Ala Gln Met Lys.Ile Arg Asp Asn Lys Ser Lys Asp Asn Ile 545 550 555 560
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- Leu Cys Pro Lys Thr Thr Tyr Ala Leu Trp Asp Leu Arg Ile Ser Leu 610 615 620
- Glu Asp Asp Val Asp Gly Glu Asp Lys Ser Ser Ser Glu Glu Ile Ser 625 630 635
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- Asp Pro Gly Phe Thr Ala Ala Pro Asn Cys Leu Ile Cys Glu Glu Leu 660 665 670
- Val Lys Thr Leu Glu Lys Arg Met Gly Lys His Pro Thr Arg Asp Ser 675 680 685
- Ile Lys His Ile Leu Glu Glu Ser Cys Asp Arg Met Arg Lys Pro Met 690 695 700
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- Asp Leu Leu Lys Glu Met Asp Pro Lys Leu Ile Cys Thr Glu Leu 725 730 735
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- Lys Gln Cys Asp Thr Phe Val Asp Gly Tyr Ala Ser Ala Val Leu Lys 820 825 830
- Leu Leu Ser Asp Val Pro Pro Lys Gln Val Cys Gln Lys Leu Gln Leu 835 840 845
- Cys Phe Ser Val Ala Val Thr Asp Glu Val Leu Glu Cys Gly Val Cys 850 860
- His Gly Val Thr Gln Ala Leu Leu Pro Phe Leu Arg Glu Lys Lys Asp 865 870 875 886
- Asn Val Ser Glu Val Thr Ala Leu Gln Met Thr Ser Val Gly Cys Glu 885 890 895
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